

SEQUENCE LISTING

<110> McCutchen, Billy F.
Herrmann, Rafael

<120> SCORPION TOXINS

<130> BB1208PCT

<140> 09/807,248

<141>

<150> 60/105,404

<151> 1998-10-23

<160> 17

<170> Microsoft Office 97

<210> 1

<211> 228

<212> DNA

<213> Leiurus quinquestriatus

<400> 1

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gtacgggtgg ccattgcgga tttaaacttg gacacggaat tgcttgcctgg tgcaatgcct 180
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<210> 2

<211> 75

<212> PRT

<213> Leiurus quinquestriatus

<220>

<221> SIGNAL

<222> (1)..(11)

<400> 2

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Leu Ala Leu Leu Phe Met Thr Gly Val Glu Ser Val Arg Asp Gly Tyr
  1             5             10             15
```

```
Ile Ala Gln Pro Glu Asn Cys Val Tyr His Cys Ile Pro Asp Cys Asp
      20             25             30
```

```
Thr Leu Cys Lys Asp Asn Gly Gly Thr Gly Gly His Cys Gly Phe Lys
      35             40             45
```

```
Leu Gly His Gly Ile Ala Cys Trp Cys Asn Ala Leu Pro Asp Asn Val
      50             55             60
```

```
Gly Ile Ile Val Asp Gly Val Lys Cys His Lys
      65             70             75
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<210> 3

<211> 238

<212> DNA

<213> Leiurus quinquestriatus

<220>

<221> unsure
 <222> (28)
 <223> n = a, c, g or t

<400> 3
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 ggaaaacggg ggtacgggtg gccattgcgg atttaaagtt ggacatggaa ctgcctgctg 180
 gtgcaatgcc ttgcccgata aagtagggat tatagtagat ggagtaaaat gccatcgc 238

<210> 4
 <211> 79
 <212> PRT
 <213> *Leiurus quinquestriatus*

<220>
 <221> SIGNAL
 <222> (1)..(12)

<400> 4
 Ser Leu Ala Leu Leu Phe Met Thr Gly Val Glu Ser Val Arg Asp Gly
 1 5 10 15
 Tyr Ile Ala Lys Pro Glu Asn Cys Ala His His Cys Phe Pro Gly Ser
 20 25 30
 Ser Gly Cys Asp Thr Leu Cys Lys Glu Asn Gly Gly Thr Gly Gly His
 35 40 45
 Cys Gly Phe Lys Val Gly His Gly Thr Ala Cys Trp Cys Asn Ala Leu
 50 55 60
 Pro Asp Lys Val Gly Ile Ile Val Asp Gly Val Lys Cys His Arg
 65 70 75

<210> 5
 <211> 258
 <212> DNA
 <213> *Leiurus quinquestriatus*

<400> 5
 atgaatcatt tggtaatgat tagtttggca cttcttttca tgacagggtg ggagagtgg 60
 gtacgtgatg ggtatattgc ccagcccga aactgtgtct accattgctt tccagggtcc 120
 cccggttgcg acacattatg taaagagaac ggtgcttcga gtggccattg cggatttaaa 180
 gaaggacacg gacttgctg ctggtgcaat gatctgcccg ataaagtagg gataatagta 240
 gaaggagaaa aatgccat 258

<210> 6
 <211> 87
 <212> PRT
 <213> *Leiurus quinquestriatus*

<220>
 <221> SIGNAL
 <222> (1)..(19)

<400> 6
 Met Asn His Leu Val Met Ile Ser Leu Ala Leu Leu Phe Met Thr Gly
 1 5 10 15
 Val Glu Ser Gly Val Arg Asp Gly Tyr Ile Ala Gln Pro Glu Asn Cys

[illegible]

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<210> 7
<211> 85
<212> PRT
<213> Buthus occitanus
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```

<400> 7
Met Ser Ser Leu Met Ile Ser Thr Ala Met Lys Gly Lys Ala Pro Tyr
  1                      5                      10                      15

Arg Gln Val Arg Asp Gly Tyr Ile Ala Gln Pro His Asn Cys Ala Tyr
          20                      25                      30

His Cys Leu Lys Ile Ser Ser Gly Cys Asp Thr Leu Cys Lys Glu Asn
          35                      40                      45

Gly Ala Thr Ser Gly His Cys Gly His Lys Ser Gly His Gly Ser Ala
          50                      55                      60

Cys Trp Cys Lys Asp Leu Pro Asp Lys Val Gly Ile Ile Val His Gly
  65                      70                      75                      80

Glu Lys Cys His Arg
          85

```

```
<210> 8
<211> 252
<212> DNA
<213> Leiurus quinquestriatus
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<220>
<221> unsure
<222> (16)
<223> n = a, c, g or t
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<400> 8
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tgcaacgatt  tatgtaccaa  gaacgggtgct  aagagtggct  attgccaatg  gttcggttca  180
agtggaaacg  cctgctgggtg  catagatttg  cccgataacg  taccgattaa  agtaccagga  240
aaatgccatc  qc                                     252
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<210> 9
<211> 84
<212> PRT
<213> Leiurus quinquestriatus
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 $\langle 220 \rangle$

<221> SIGNAL
<222> (1)..(19)

<220>
<221> UNSURE
<222> (6)
<223> Xaa = ANY AMINO ACID

<400> 9
Met Asn Tyr Leu Val Xaa Ile Ser Leu Ala Leu Leu Leu Met Thr Gly
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Val Glu Ser Gly Arg Asp Ala Tyr Ile Ala Gln Asn Tyr Asn Cys Val
20 25 30
Tyr His Cys Ala Leu Asn Pro Tyr Cys Asn Asp Leu Cys Thr Lys Asn
35 40 45
Gly Ala Lys Ser Gly Tyr Cys Gln Trp Phe Gly Ser Ser Gly Asn Ala
50 55 60
Cys Trp Cys Ile Asp Leu Pro Asp Asn Val Pro Ile Lys Val Pro Gly
65 70 75 80
Lys Cys His Arg

<210> 10
<211> 65
<212> PRT
<213> Buthus occitanus tunetanus

<400> 10
Gly Arg Asp Ala Tyr Ile Ala Gln Pro Glu Asn Cys Val Tyr Glu Cys
1 5 10 15
Ala Gln Asn Ser Tyr Cys Asn Asp Leu Cys Thr Lys Asn Gly Ala Thr
20 25 30
Ser Gly Tyr Cys Gln Trp Leu Gly Lys Tyr Gly Asn Ala Cys Trp Cys
35 40 45
Lys Asp Leu Pro Asp Asn Val Pro Ile Arg Ile Pro Gly Lys Cys His
50 55 60
Phe
65

<210> 11
<211> 256
<212> DNA
<213> Leiurus quinquestriatus

<400> 11
atgaaactct tactttttact cattgtctct gcttcaatgc tgattgaaag cttagttaat 60
gctgacggat atataagaag aaaagacgga tgcaagggtg catgcctgtt cggaaatgac 120
ggctgcaata aagaatgcaa agcttatggt gcctattatg gatattgttg gacctgggga 180
cttgctgct ggtgcgaagg tcttcggat gacaagacat ggaagagtga aacaaacaca 240
tgcggtggca aaaagt 256

<210> 12

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<400> 14
Met Lys Leu Leu Leu Leu Leu Thr Ile Ser Ala Ser Met Leu Ile Glu
  1                      5                      10                      15

Gly Leu Val Asn Ala Asp Gly Tyr Ile Arg Gly Gly Asp Gly Cys Lys
          20                      25                      30

Val Ser Cys Val Ile Asn His Val Phe Cys Asp Asn Glu Cys Lys Ala
      35                      40                      45

Ala Gly Gly Ser Tyr Gly Tyr Cys Trp Ala Trp Gly Leu Ala Cys Trp
  50                      55                      60

```

Cys Glu Gly Leu Pro Ala Asp Arg Glu Trp Lys Tyr Glu Thr Asn Thr
 65 70 75 80

Cys Gly Gly Lys Lys
 85

<210> 15
 <211> 255
 <212> DNA
 <213> Leiurus quinquestriatus

<400> 15
 atgaaaataa taatttttct aattgtgtca tcattaatgc tgataggagt gaagaccgat 60
 aatgggttact tgcttaacaa agccaccggt tgcaaggtct ggtgtgttat taataatgca 120
 tcttgtaata gtgagtgtaa actaagacgt ggaaattatg gctactgcta tttctggaaa 180
 ttggcctgtt attgcgaagg agctccaaaa tcagaacttt gggcttacgc aaccaataaa 240
 tgcaatggga aatta 255

<210> 16
 <211> 85
 <212> PRT
 <213> Leiurus quinquestriatus

<220>
 <221> SIGNAL
 <222> (1)..(19)

<400> 16
 Met Lys Leu Leu Leu Leu Leu Ile Val Ser Ala Ser Met Leu Ile Glu
 1 5 10 15

Ser Leu Val Asn Ala Asp Gly Tyr Ile Arg Arg Lys Asp Gly Cys Lys
 20 25 30

Val Ala Cys Leu Phe Gly Asn Asp Gly Cys Asn Lys Glu Cys Lys Ala
 35 40 45

Tyr Gly Ala Tyr Tyr Gly Tyr Cys Trp Thr Trp Gly Leu Ala Cys Trp
 50 55 60

Cys Glu Gly Leu Pro Asp Asp Lys Thr Trp Lys Ser Glu Thr Asn Thr
 65 70 75 80

Cys Gly Gly Lys Lys
 85

<210> 17
 <211> 61
 <212> PRT
 <213> Leiurus quinquestriatus

<400> 17
 Asp Gly Tyr Ile Lys Arg Arg Asp Gly Cys Lys Val Ala Cys Leu Ile
 1 5 10 15

Gly Asn Glu Gly Cys Asp Lys Glu Cys Lys Ala Tyr Gly Gly Ser Tyr
 20 25 30

Gly Tyr Cys Trp Thr Trp Gly Leu Ala Cys Trp Cys Glu Gly Leu Pro
 35 40 45

Asp Asp Lys Thr Trp Lys Ser Glu Thr Asn Thr Cys Glu
50 55 60